
Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=8; day=27; hr=11; min=54; sec=57; ms=562;]

Validated By CRFValidator v 1.0.3

Application No: 10534081 Version No: 1.0

Input Set:

Output Set:

Started: 2008-08-23 06:09:18.527 **Finished:** 2008-08-23 06:09:20.489

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 962 ms

Total Warnings: 16
Total Errors: 0

No. of SeqIDs Defined: 26

Actual SeqID Count: 26

Error code		Error Description											
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(11)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(12)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(13)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(14)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(15)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(16)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(17)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(18)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(19)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(20)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(21)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(22)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(23)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(24)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(25)		
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(26)		

SEQUENCE LISTING

```
<110> FUKATSU, KOHJI
     SASAKI, SHINOBU
     HINUMA, SHUJI
     ITO, YASUAKI
     SUZUKI, NOBUHIRO
     HARADA, MASATAKA
     YASUMA, TSUNEO
<120> RECEPTOR FUNCTION REGULATOR
<130> 66530 (46590)
<140> 10534081
<141> 2008-08-23
<150> PCT/JP2003/014139
<151> 2003-11-06
<150> JP 2003-153986
<151> 2003-05-30
<150> JP 2003-16889
<151> 2003-01-27
<150> JP 2002-324632
<151> 2002-11-08
<160> 26
<170> PatentIn Ver. 3.3
<210> 1
<211> 300
<212> PRT
<213> Mus musculus
<400> 1
Met Asp Leu Pro Pro Gln Leu Ser Phe Ala Leu Tyr Val Ser Ala Phe
                                   10
Ala Leu Gly Phe Pro Leu Asn Leu Leu Ala Ile Arg Gly Ala Val Ser
                               25
His Ala Lys Leu Arg Leu Thr Pro Ser Leu Val Tyr Thr Leu His Leu
                           40
Gly Cys Ser Asp Leu Leu Leu Ala Ile Thr Leu Pro Leu Lys Ala Val
    50 55 60
Glu Ala Leu Ala Ser Gly Ala Trp Pro Leu Pro Leu Pro Phe Cys Pro
                    70
65
                                      75
```

Val Phe Ala Leu Ala His Phe Ala Pro Leu Tyr Ala Gly Gly Phe

90

95

85

Leu Ala Ala Leu Ser Ala Gly Arg Tyr Leu Gly Ala Ala Phe Pro Phe 100 105 110

Gly Tyr Gln Ala Ile Arg Arg Pro Arg Tyr Ser Trp Gly Val Cys Val
115 120 125

Ala Ile Trp Ala Leu Val Leu Cys His Leu Gly Leu Ala Leu Gly Leu 130 135 140

Glu Thr Ser Gly Ser Trp Leu Asp Asn Ser Thr Ser Ser Leu Gly Ile 145 150 155 160

Asn Ile Pro Val Asn Gly Ser Pro Val Cys Leu Glu Ala Trp Asp Pro 165 170 175

Asp Ser Ala Arg Pro Ala Arg Leu Ser Phe Ser Ile Leu Leu Phe Phe 180 185 190

Leu Pro Leu Val Ile Thr Ala Phe Cys Tyr Val Gly Cys Leu Arg Ala 195 200 205

Leu Val Arg Ser Gly Leu Ser His Lys Arg Lys Leu Arg Ala Ala Trp
210 215 220

Val Ala Gly Gly Ala Leu Leu Thr Leu Leu Cys Leu Gly Pro Tyr 225 230 235 240

Asn Ala Ser Asn Val Ala Ser Phe Ile Asn Pro Asp Leu Gly Gly Ser 245 250 255

Trp Arg Lys Leu Gly Leu Ile Thr Gly Ala Trp Ser Val Val Leu Asn 260 265 270

Pro Leu Val Thr Gly Tyr Leu Gly Thr Gly Pro Gly Arg Gly Thr Ile 275 280 285

Cys Val Thr Arg Thr Gln Arg Gly Thr Ile Gln Lys 290 295 300

<210> 2

<211> 900

<212> DNA

<213> Mus musculus

<400> 2

atggacctge ceceacaget ctectteget etetatgtat etgeetttge getgggettt 60 ceattgaact tgttagecat eegaggegea gtgteecaeg etaaactgeg acteactee 120 agettggtet acactetee tetgggetge tetgatetee taetggeeat eactetgee 240 gtetttgeet tggaegeet tgeeceete taeegaggeg gaggetteet agetgetee 30 agegetggee getacetggg ggetgeette eecttegggt accaageeat eeggaggeee 36 egetatteet ggggtgtgt tgtggetata tgggeeettg teetetgea eetggggetgeet eeggaggee 420 geetttgget tggaagetg eeggaagetg eeggaageetg eeggaggeetg teggaageetg eeggaggeetg eeggaageetg eeggaage

tgctatgtgg gctgcctccg ggccctggtg cgctcaggcc tgagccacaa acggaagctc 660 agggcagctt gggtggccgg aggcgctctc ctcacactcc tgctctgcct ggggccctat 720 aatgcctcca atgtggctag tttcataaac ccggacctag gaggctcctg gaggaagttg 780 ggactcatca caggggcctg gagtgtggta ctcaacccac tggtcactgg ctacttggga 840 acaggtcctg gacgggaac aatatgtgtg acgaggactc aaagaggaac aattcagaag 900

<210> 3

<211> 300

<212> PRT

<213> Rattus norvegicus

<400> 3

Met Asp Leu Pro Pro Gln Leu Ser Phe Ala Leu Tyr Val Ser Ala Phe 1 5 10 15

Ala Leu Gly Phe Pro Leu Asn Leu Leu Ala Ile Arg Gly Ala Val Ser 20 25 30

His Ala Lys Leu Arg Leu Thr Pro Ser Leu Val Tyr Thr Leu His Leu 35 40 45

Ala Cys Ser Asp Leu Leu Leu Ala Ile Thr Leu Pro Leu Lys Ala Val 50 55 60

Glu Ala Leu Ala Ser Gly Val Trp Pro Leu Pro Leu Pro Phe Cys Pro
65 70 75 80

Val Phe Ala Leu Ala His Phe Ala Pro Leu Tyr Ala Gly Gly Gly Phe
85 90 95

Leu Ala Ala Leu Ser Ala Gly Arg Tyr Leu Gly Ala Ala Phe Pro Phe $100 \,$ $105 \,$ $110 \,$

Gly Tyr Gln Ala Ile Arg Arg Pro Cys Tyr Ser Trp Gly Val Cys Val
115 120 125

Ala Ile Trp Ala Leu Val Leu Cys His Leu Gly Leu Ala Leu Gly Leu 130 135 140

Glu Ala Pro Arg Gly Trp Val Asp Asn Thr Thr Ser Ser Leu Gly Ile 145 150 155 160

Asn Ile Pro Val Asn Gly Ser Pro Val Cys Leu Glu Ala Trp Asp Pro 165 170 175

Asp Ser Ala Arg Pro Ala Arg Leu Ser Phe Ser Ile Leu Leu Phe Phe 180 185 190

Leu Pro Leu Val Ile Thr Ala Phe Cys Tyr Val Gly Cys Leu Arg Ala 195 200 205

Leu Val His Ser Gly Leu Ser His Lys Arg Lys Leu Arg Ala Ala Trp 210 215 220

Val Ala Gly Gly Ala Leu Leu Thr Leu Leu Cys Leu Gly Pro Tyr 225 230 235 240

```
Asn Ala Ser Asn Val Ala Ser Phe Ile Asn Pro Asp Leu Glu Gly Ser
                245
                                                         255
Trp Arg Lys Leu Gly Leu Ile Thr Gly Ala Trp Ser Val Val Leu Asn
            260
                                2.65
                                                    270
Pro Leu Val Thr Gly Tyr Leu Gly Thr Gly Pro Gly Gln Gly Thr Ile
                            280
Cys Val Thr Arg Thr Pro Arg Gly Thr Ile Gln Lys
                        295
    290
<210> 4
<211> 900
<212> DNA
<213> Rattus norvegicus
<400> 4
atggacctgc ccccacagct ctccttcgct ctctatgtat cagcctttgc actaggcttt 60
ccattgaact tgttagccat ccgaggtgca gtgtcccacg cgaaactgcg actcaccccc 120
agettggtct acacteteca tttggcetge tetgacetee tactggeeat caecetgeee 180
ctgaaggctg tggaggccct ggcttctggg gtctggcccc tgccactccc cttctgccca 240
gtctttgcct tggcccactt tgcgcccctc tatgcaggtg gaggcttcct ggctgctctc 300
agtgctggcc gctacctggg agctgccttc ccctttggat accaagccat ccggaggccc 360
tgctattcct ggggtgtgtg tgtggctata tgggcccttg tcctttgcca cctgggactg 420
getettgget tggaggetee eagaggetgg gtggataaca ceaceagtte eetgggeate 480
aacatacccg tgaatggctc cccggtctgc ctggaagcgt gggatcctga ctctgcccgc 540\,
cctgcccgac tcagtttctc gattctgctc ttctttctgc ccttggttat cactgctttc 600
tgctatgtgg gctgcctccg ggccctggtg cactcgggcc tgagccacaa acggaagctc 660
agggcagett gggtggetgg aggageaett eteacaetee tgetetgeet ggggeeetat 720
aatgcttcca atgtggctag tttcataaac ccggacttag aaggctcctg gaggaagttg 780
gggctcatca caggagcctg gagtgtggtg ctcaacccac tggtcactgg ctacttggga 840
acaggtcctg gacaggggac aatatgtgtg accaggactc caagagggac aattcagaag 900
<210> 5
<211> 300
<212> PRT
<213> Homo sapiens
<400> 5
Met Asp Leu Pro Pro Gln Leu Ser Phe Gly Leu Tyr Val Ala Ala Phe
Ala Leu Gly Phe Pro Leu Asn Val Leu Ala Ile Arg Gly Ala Thr Ala
             2.0
                                 25
His Ala Arg Leu Arg Leu Thr Pro Ser Leu Val Tyr Ala Leu Asn Leu
         35
Gly Cys Ser Asp Leu Leu Thr Val Ser Leu Pro Leu Lys Ala Val
     50
                         55
Glu Ala Leu Ala Ser Gly Ala Trp Pro Leu Pro Ala Ser Leu Cys Pro
```

65

70

75

Val Phe Ala Val Ala His Phe Phe Pro Leu Tyr Ala Gly Gly Phe Leu Ala Ala Leu Ser Ala Gly Arg Tyr Leu Gly Ala Ala Phe Pro Leu 105 100

110

Gly Tyr Gln Ala Phe Arg Arg Pro Cys Tyr Ser Trp Gly Val Cys Ala 120

Ala Ile Trp Ala Leu Val Leu Cys His Leu Gly Leu Val Phe Gly Leu 130 135 140

Glu Ala Pro Gly Gly Trp Leu Asp His Ser Asn Thr Ser Leu Gly Ile 150 155

Asn Thr Pro Val Asn Gly Ser Pro Val Cys Leu Glu Ala Trp Asp Pro 165 170

Ala Ser Ala Gly Pro Ala Arg Phe Ser Leu Ser Leu Leu Leu Phe Phe 180 185

Leu Pro Leu Ala Ile Thr Ala Phe Cys Tyr Val Gly Cys Leu Arg Ala 195 200

Leu Ala Arg Ser Gly Leu Thr His Arg Arg Lys Leu Arg Ala Ala Trp 210 215 220

Val Ala Gly Gly Ala Leu Leu Thr Leu Leu Cys Val Gly Pro Tyr 230 235

Asn Ala Ser Asn Val Ala Ser Phe Leu Tyr Pro Asn Leu Gly Gly Ser 245 250

Trp Arg Lys Leu Gly Leu Ile Thr Gly Ala Trp Ser Val Val Leu Asn 260

Pro Leu Val Thr Gly Tyr Leu Gly Arg Gly Pro Gly Leu Lys Thr Val 275 280

Cys Ala Ala Arg Thr Gln Gly Gly Lys Ser Gln Lys 290 295 300

<210> 6 <211> 900

<212> DNA

<213> Homo sapiens

<400> 6

atggacctgc ccccqcaqct ctccttcgqc ctctatgtgq ccqcctttqc gctqgqcttc 60 ccgctcaacg tcctggccat ccgaggcgcg acggcccacg cccggctccg tctcacccct 120 agcctggtct acgccctgaa cctgggctgc tccgacctgc tgctgacagt ctctctgccc 180 ctgaaggcgg tggaggcgct agcctccggg gcctggcctc tgccggcctc gctgtgcccc 240 gtettegegg tggcceaett etteceaete tatgeeggeg ggggetteet ggeegeeetg 300 agtgcaggcc gctacctggg agcagccttc cccttgggct accaagcctt ccggaggccg 360 tgctattcct ggggggtgtg cgcggccatc tgggccctcg tcctgtgtca cctgggtctq 420 gtetttgggt tggaggete aggaggetgg etggaecaca geaacacete eetgggeate 480 aacacacegg teaacggete teeggtetge etggaggeet gggaecegge etetgeegge 540 eeggeecget teageetete teteetgete tttttetge eettggeeat eacageette 600 tgetaegtgg getgeeteeg ggeactggee egeteeggee tgaegeacag geggaagetg 660 egggeegeet gggtggeegg eggggeeete eteaegetge tgetetgegt aggaecetae 720 aacgeeteea aegtggeeag etteetgtae eecaatetag gaggeteetg geggaagetg 780 gggeteatea egggtgeetg gagtgtggt ettaateege tggtgaecgg ttaettggga 840 aggggteetg geetgaagae agtgtgtgeg geaagaacge aaggggeaa gteecagaag 900

<210> 7

<211> 300

<212> PRT

<213> Macaca fascicularis

<400> 7

Met Asp Leu Pro Pro Gln Leu Ser Phe Ala Leu Tyr Val Ala Ala Phe 1 5 10 15

Ala Leu Gly Phe Pro Leu Asn Val Leu Ala Ile Arg Gly Ala Arg Ala
20 25 30

His Ala Arg Arg Leu Thr Pro Ser Leu Val Tyr Ala Leu Asn Leu 35 40 45

Gly Cys Ser Asp Leu Leu Leu Thr Val Ser Leu Pro Leu Lys Ala Val
50 55 60

Glu Ala Leu Ala Ser Gly Ala Trp Pro Leu Pro Ala Ser Leu Cys Pro
65 70 75 80

Val Phe Gly Val Ala His Phe Ala Pro Leu Tyr Ala Gly Gly Gly Phe
85 90 95

Leu Ala Ala Leu Ser Ala Gly Arg Tyr Leu Gly Ala Ala Phe Pro Leu 100 105 110

Gly Tyr Gln Ala Phe Arg Arg Pro Cys Tyr Ser Trp Gly Val Cys Ala 115 120 125

Ala Ile Trp Ala Leu Val Leu Cys His Leu Gly Leu Val Phe Val Leu 130 135 140

Glu Ala Pro Gly Gly Trp Leu Asp His Ser Asn Thr Ser Leu Gly Ile 145 150 155 160

Asn Thr Pro Val Asn Gly Ser Pro Val Cys Leu Glu Ala Trp Asp Pro 165 170 175

Ala Ser Ala Gly Pro Ala Arg Phe Ser Leu Ser Leu Leu Leu Phe Phe 180 185 190

Leu Pro Leu Ala Ile Thr Ala Phe Cys Tyr Val Gly Cys Leu Arg Ala 195 200 205

Leu Ala His Ser Gly Leu Thr His Arg Arg Lys Leu Arg Ala Ala Trp
210 215 220

```
Val Ala Gly Gly Ala Leu Leu Thr Leu Leu Leu Cys Val Gly Pro Tyr
225
                    230
                                         235
                                                              240
Asn Ala Ser Asn Val Ala Ser Phe Leu Asn Pro Asn Leu Gly Gly Ser
                245
                                     250
                                                          2.55
Trp Arg Lys Leu Gly Leu Ile Thr Gly Ala Trp Ser Val Val Leu Asn
            260
                                 265
Pro Leu Val Thr Gly Tyr Leu Gly Arg Gly Pro Gly Leu Lys Thr Val
                             280
        275
                                                  285
Cys Ala Ala Arg Thr Gln Gly Ser Thr Ser Gln Lys
                        295
<210> 8
<211> 900
<212> DNA
<213> Macaca fascicularis
<400> 8
atggacctgc ccccgcagct ctcctttgcc ctctatgtgg cggcctttgc gctgggcttc 60
ccgctcaacg tcctggccat ccgaggggcg agggcccacg cccggcgccg tctcaccccc 120
agectggtet aegecetgaa eetgggetge teegaeetgt tgetgaeagt eteeetgeee 180
\verb|ctgaaggcgg|| \verb|tggaggcgct|| \verb|gcctccggg|| \verb|gcctggcctc|| tgccggcctc|| actgtgccct|| 240
gtcttcgggg tggcccactt tgctccactc tatgccggcg ggggcttcct ggccgccctg 300
agtgcaggcc gctacctggg agcggccttc cccttgggct accaagcctt ccggaggccg 360
tgctattcct ggggggtgtg tgcggccatc tgggccctcg tcctgtgtca cctgggtctg 420
gtctttgtgt tggaggctcc gggaggctgg ctggaccaca gcaacacctc actgggcatc 480
aacacaccgg tcaacggctc tcccgtctgc ctggaggcct gggacccggc ctctgccggc 540
ccggcccgct tcagcctctc tctcctgctt tttttcctgc ccttggccat cacagccttc 600
tgctacgtgg gctgcctccg ggcactggcc cactccggcc tgacccacag gcggaagctg 660
agggccgcct gggtagccgg cggggccctc ctcacgctgc tgctctgcgt aggaccctac 720
aacgcctcca atgtggccag ctttctgaac cccaatctgg gaggctcctg gcggaagctg 780
gggctcatca cgggtgcctg gagtgtggtg ctcaacccgc tggtgaccgg ttacttggga 840
aggggtcctg gcctgaagac agtgtgtgcg gcaagaacgc aagggagcac gtcccagaag 900
<210> 9
<211> 300
<212> PRT
<213> Mesocricetus auratus
<400> 9
Met Ala Leu Ser Pro Gln Leu Phe Phe Ala Leu Tyr Val Ser Ala Phe
                                      1.0
Ala Leu Gly Phe Pro Leu Asn Leu Leu Ala Ile Arg Gly Ala Val Ala
             20
                                  25
Arg Ala Arg Leu Arg Leu Thr Pro Asn Leu Val Tyr Thr Leu His Leu
         35
                              40
```

Ala Cys Ser Asp Leu Leu Ala Ile Thr Leu Pro Val Lys Ala Val

60

55

50

Glu 65	Ala	Leu	Ala	Ser	Gly 70	Ala	Trp	Pro	Leu	Pro 75	Leu	Pro	Leu	Суз	Pro 80
Val	Phe	Val	Leu	Val 85	His	Phe	Ala	Pro	Leu 90	Tyr	Ala	Gly	Gly	Gly 95	Phe
Leu	Ala	Ala	Leu 100	Ser	Ala	Gly	Arg	Tyr 105	Leu	Gly	Ala	Ala	Phe 110	Pro	Phe
Gly	Tyr	Gln 115	Ala	Val	Arg	Arg	Pro 120	Arg	Tyr	Ser	Trp	Gly 125	Val	Cys	Val
Ala	Ile 130	Trp	Ala	Leu	Val	Leu 135	Cys	His	Met	Gly	Leu 140	Val	Leu	Gly	Leu
Glu 145	Ala	Pro	Gly	Gly	Trp 150	Leu	Asn	Thr	Thr	Ser 155	Ser	Ser	Leu	Gly	Ile 160
Asn	Thr	Pro	Val	Asn 165	Gly	Ser	Pro	Val	Cys 170	Leu	Glu	Ala	Trp	Asp 175	Pro
Asn	Ser	Ala	Arg 180	Pro	Ala	Arg	Leu	Ser 185	Phe	Ser	Ile	Leu	Leu 190	Phe	Phe
Val	Pro	Leu 195	Val	Ile	Thr	Ala	Phe 200	Суз	Tyr	Val	Gly	Cys 205	Leu	Arg	Ala
Leu	Ala 210	His	Ser	Gly	Leu	Ser 215	His	Lys	Arg	Lys	Leu 220	Arg	Ala	Ala	Trp
Ala 225	Ala	Gly	Gly	Ala	Phe 230	Leu	Thr	Leu	Leu	Leu 235	Cys	Leu	Gly	Pro	Tyr 240
Asn	Ala	Ser	Asn	Val 245	Ala	Ser	Phe	Val	Asn 250	Pro	Asp	Leu	Gly	Gly 255	Ser
Trp	Arg	Lys	Leu 260	Gly	Leu	Ile	Thr	Gly	Ser	Trp	Ser	Val	Val	Leu	Asn